

Region	AG1 (0-10 years)	AG2 (10-30 years)	AG3 (30-40 years)	AG4 (40-50 years)	AG5 (50-60 years)	AG6 (60-above years)	References
American (AM)	NA	NA	NA	NA	NA	NA	1
Danish (DA)	NA	NA	NA				2
Spanish (ES)	NA						2
French (FR)	NA	NA	NA	NA	NA		3
Italian (IT)	NA	NA	NA	NA	NA		3
Chinese (CH)	NA						4
Indian (IN)		NA	NA	NA	NA	NA	5
Japanese (JP)					NA	NA	3

**References:**

1. Human Microbiome Project Consortium. Structure, function and diversity of the healthy human microbiome. *Nature*. 2012;486: 207-214.
2. Qin J, Li R, Raes J, Arumugam M, Burgdorf KS, Manichanh C et al. A human gut microbial gene catalogue established by metagenomic sequencing. *Nature*. 2010;464: 59-65.
3. Arumugam M, Raes J, Pelletier E, Le Paslier D, Yamada T, Mende DR et al. Enterotypes of the human gut microbiome. *Nature*. 2011;473: 174-180.
4. Qin J, Li Y, Cai Z, Li S, Zhu J, Zhang F et al. A metagenome-wide association study of gut microbiota in type 2 diabetes. *Nature*. 2012;490: 55-60.
5. Ghosh TS, Gupta SS, Bhattacharya T, Yadav D, Barik A, Chowdhury A et al. Gut microbiomes of Indian children of varying nutritional status. *PLoS One*. 2014;9: e95547.

